OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/052,283

DATE: 02/12/2002
TIME: 12:01:23

Input Set : A:\P2751R1C1.txt
Output Set: N:\CRF3\02122002\J052283.raw

p5

ENTERED

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 3 <110> APPLICANT: Baker,Kevin P.
 4        Goddard,Audrey
 5        Wood,William I.
 7 <120> TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES AND METHOD OF USE THEREOF
 9 <130> FILE REFERENCE: P2751R1C1
11 <140> CURRENT APPLICATION NUMBER: US/10/052,283
11 <141> CURRENT FILING DATE: 2002-01-15
11 <150> PRIOR APPLICATION NUMBER: PCT/US00/20006
12 <151> PRIOR FILING DATE: 2000-07-21
14 <150> PRIOR APPLICATION NUMBER: US 60/145,701
15 <151> PRIOR FILING DATE  1999-07-26
17 <160> NUMBER OF SEQ ID NOS: 564
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21 <212> TYPE: DNA
22 <213> ORGANISM: Homo Sapien
24 <220> FEATURE:
25 <221> NAME/KEY: unsure
26 <222> LOCATION: 212, 285, 424
27 <223> OTHER INFORMATION  unknown base
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32  tatacagatg cacgcttatt tatacttatg tgtaagtgaa ataaatggca 100
34  aaaatgatac aaggcatagg aagaagaaat taggattata tgctatgtaa 150
36  gaagcagtat agtgtttttt gaaaatagac ttgaattagt tggaaatcca 200
38  tattgaaaac tntcgggcaa acattttaa aaaataaaaa aatgatatgc 250
40  taagaaagaa gagaaacgg aattacacaa aatgntcaat taaaaccaca 300
42  aaaggaagca aaagtgtgga aaacaaaag gggaacaaag aataaggcaa 350
44  caaacagaaa acagtaacaa atatggtaag cattaatcca actatattaa 400
46  taatcacttt aaatatcaat ggtntaaata tgtcaattat aagacagaga 450
48  ttaccagagt ggacacatta tataagct 478
50 <210> SEQ ID NO: 2
51 <211> LENGTH: 682
52 <212> TYPE: DNA
53 <213> ORGANISM: Homo Sapien
55 <220> FEATURE:
56 <221> NAME/KEY: unsure
57 <222> LOCATION: 45, 51, 62
58 <223> OTHER INFORMATION: unknown base
60 <400> SEQUENCE: 2
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63  nttcccgccc gnttccggcc ccccagtgcc gcccttcgcg ggccggaggc 100
65  gcgagtctgg gctttggcgc cttcgcagcc gcaggcgaca tcctctttcc 150
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W--> 38
W--> 40
W--> 46
W--> 61
W--> 63

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 67    tagctaaagc cccaaacgcc caggtggctt cctgggagag cacggctgag 200
 69    cctccgcctt cagatcagaa caggcagagc ctccaagggc ggcttgggcc 250
 71    cagtgcctgc ttatcctgcc cgtctctcca cacttcttct ttccctgtcc 300
 73    gttggagtcc attccttcct ggaaaagcca aagccgcgct cccctaagag 350
 75    tcatgtgtta ctggattaat tgaaattctt gataggtaac agagttttat 400
 77    catcagctta tgattgccta tgactagctc aaagttaagt tttaataaac 450
 79    tagtaagtac aataaacctc attcctaaat acaaggaaaa gaactatata 500
 81    atgaatactt gtctctatgc ccctctcgca tagataacaa taatttaggt 550
 83    ttacctttaa atgaactgca tttaaatgaa attaatttaa atgattgttt 600
 85    cacggcacag tttcatcaat ggtctacggt atcccttatt tatgtataca 650
 87    tcagtttgta tacatctgta tctatgtatt tg 682
 89  <210> SEQ ID NO: 3
 90  <211> LENGTH: 835
 91  <212> TYPE: DNA
 92  <213> ORGANISM: Homo Sapien
 94  <220> FEATURE:
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 96  <222> LOCATION: 1, 6, 12, 27, 29-31, 49, 53, 56, 58, 61-62, 66, 68, 71, 74,
 97        78-81, 85-86, 88, 91, 95, 98, 102, 105, 109, 111, 115, 126-127,
 98        131, 139, 142, 146-147, 157, 163, 166, 169, 175, 177, 184,
 99        196-197, 199, 210, 215, 217, 219, 222, 228, 234, 236, 238,
100        240-241, 244-245, 257, 260, 263, 271, 279, 287, 291-331, 335,
101        337, 345, 384, 392, 418, 438, 459, 465, 478, 499, 501, 507, 533,
102        562, 579, 587, 598, 616, 630, 656, 661, 667, 671, 684, 702, 746,
103        811, 824-835
104  <223> OTHER INFORMATION: unknown base
106  <400> SEQUENCE: 3
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W--> 109    ganggngnag nngagnangg nggnggannn naggnngnag naggnccnag 100
W--> 111    anggnaagng nttgnaagaa agggnanngcc nggtaaaana gnaccnnccc 150
W--> 113    aagaagngat tanggnggnt tcctngntga aggntgtgga tcccanntnt 200
W--> 115    ttcccgggan ttatngntng gnaacaanat ttcnangngn nacnnaggca 250
W--> 117    aacaatnaan ttnccaaggt nttggtagna tttcccncgg nnnnnnnnnn 300
W--> 119    nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn ngatntnggg gtttntcccc 350
W--> 121    tttcccttc cccttccccc accccggggg ttcnggttgg tnaagaaaaa 400
W--> 123    aaaaaaaaaa aagaattntg gcgcggcctc ggcggagntg gtgatcggct 450
W--> 125    ggtgcatant cggcntctta ctactggnta ttttggcatt ctgctggana 500
W--> 127    natgttngta aataccaaag tcggcgggaa agngaagttg tttccaccat 550
W--> 129    aacagcaatt tnttttttag caattgcant tatcacntca gcacttgnac 600
W--> 131    cagaggaaat attttnggtt tcttacatgn aaaatcaaaa tggtacattt 650
W--> 133    aaggantggg ntaatgntaa ngtcagcaga cagnttgagg acactgtatt 700
W--> 135    anacggttac tatactttat attctgttat attgttctgt gtgttnttct 750
     137    ggatcccttt tgtctactta tattatgaag aaaaggatga tgatgatact 800
W--> 139    agtagatgta ntcaaattaa aacnnnnnnn nnnnn 835
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     142  <211> LENGTH: 491
     143  <212> TYPE: DNA
     144  <213> ORGANISM: Homo Sapien
     146  <220> FEATURE:
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      149        403, 427, 442, 447, 451, 463, 481, 484, 486, 489
      150  <223> OTHER INFORMATION: unknown base
      152  <400> SEQUENCE: 4
      153  agtttgttaa aaataataat gccaataata tatgttattt aacgtatgtt 50
      155  tataacagat gcacgcttat ttatacttat gtgtaagtga aataaatggc 100
      157  aaaaatgata caaggcatag gaagaagaaa ttaggattat atgctatgta 150
W-->  159  agaagcagta tagtgttttt tgaaaataga nttgaattag ttggaaatcc 200
W-->  161  atattgaaaa ctntcgggca aacattttta aaaaataaaa aaatgatatg 250
W-->  163  ntaagaaaga agagaaaacg gaattacaca aaatgctcaa ttaaaaccac 300
W-->  165  aaaaggaagc aaaagtgtgg aaaacaaaaa ggggnacaaa gaatnnggcn 350
W-->  167  acaaacngca aacagtaaca attntggtaa ncattantcc aattatantt 400
W-->  169  ncnattactc taaatatcaa tgttttnaat atgtctattg tnagacngag 450
W-->  171  nttaccagag agnacacatt atataaggtc ngangngtng g 491
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      174  <211> LENGTH: 637
      175  <212> TYPE: DNA
      176  <213> ORGANISM: Homo Sapien
      178  <220> FEATURE:
      179  <221> NAME/KEY: unsure
      180  <222> LOCATION: 4, 12-13, 31, 37, 40, 43-44, 46, 50, 55, 64, 68, 78, 97, 114,
      181        124-125, 137, 148, 183, 185, 227, 233, 236, 238, 264, 266, 268,
      182        276, 322, 325, 334, 340, 345, 417, 426, 428, 472, 474-475,
      183        482-484, 500, 502-503, 509-510, 521, 540, 543, 558, 588-589, 594,
      184        598, 606, 616, 631, 633
      185  <223> OTHER INFORMATION: unknown base
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W-->  188  ttcnttgtca anngtttttg gttccccctt ntttccnggn ttnntntttn 50
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W-->  192  ggattgaatt cccngggggat cttnnagaga tcccttngac tttgaccnaa 150
W-->  194  gggtccggct ttaggggaag aagttggtgt ttngntgggc cctggtactg 200
W-->  196  aagacgcgtt ccgggtagcc caaagangtt tcntantnac ccaaagcccc 250
W-->  198  gcacccgcct tttntntntt ttcttntggc aggatgaggc gtgcaggcct 300
W-->  200  gggtgaagga gtacttcctg gnaantatgg gaantatggn tatgntaata 350
      202  gtgggtatag tgcctgtgaa gaagaaaatg agaggctcac tgaaagtttg 400
W-->  204  agaagcaaag taactgntat aaaatntntt tcccattgaa ataggccatg 450
W-->  206  aagttaaaac ccagaataaa tnannagcgg annnggatta aaaagacgan 500
W-->  208  tnnacaacnn tgattttgta ngtataacta tgggcataan tgnagatttt 550
W-->  210  ttccagangg agctaaacaa agatgttgtg agatatgnng aggntatnat 600
W-->  212  taattntcaa gtttgntcac ataggcgagc ntnaaaac 637
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      215  <211> LENGTH  969
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      217  <213> ORGANISM: Homo Sapien
      219  <220> FEATURE:
      220  <221> NAME/KEY: unsure
      221  <222> LOCATION: 12-13, 25, 46, 49, 53, 64, 94, 118, 134, 140, 153, 155, 164, 172,
      222        175, 177, 182, 184, 191, 210, 218, 221, 223-224, 240, 242, 250,
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    Input Set   A:\P2751R1C1.txt
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      223         259, 266-267, 272, 276, 298, 305-306, 319, 339-340, 344, 346,
      224         348, 353, 384, 397, 408, 432, 455, 457, 470, 474, 480, 502, 504,
      225         519, 525, 559, 579, 597, 625, 629, 631, 634, 645, 649, 655, 657,
      226         666, 670, 675, 681, 697, 768, 773, 776, 786, 788, 791, 794, 816,
      227         819, 830, 859, 877, 879, 886, 895, 904, 910, 913, 918, 925,
      228         927-946, 952, 956, 958, 962, 966, 968-969
      229  <223> OTHER INFORMATION: unknown base
      231  <400> SEQUENCE: 6
W--> 232  cccctttttc cnnggttttt ttttnggaaa aaaatttcag gggtanccng 50
W--> 234  ggnaaaattt aaantccagg gtttgggggg atttccccgg ggtncttttg 100
W--> 236  gagttccttt ggacctgnaa caaagggttg gaantaaaan aaaaattaaa 150
W--> 238  aancngggtt tttngggggaa anttnanaat gngnttggggg ncaagaaaaa 200
W--> 240  tgggtttttn gggagggnaa ngnnggttca tttccaaatn gnagggggggn 250
W--> 242  aaaaatttna ggcttnnggg gnaggnggaa aaaaatttcg tagcctcnag 300
W--> 244  gttgnnattt ttaaacctnc agaaggtggc cagccccgnn tcancngntg 350
W--> 246  atnaaggcag atgggaaaag ggggatatgg ggtnataagg gtacctntca 400
W--> 248  cccttttnga aggaaaaaaa gtggtccaca gnatttttgt ttacccaagg 450
W--> 250  gtaananatg gaattttgtn gaanataggn gaatggtgag gcatttggaa 500
W--> 252  ananggggggg gggttttnt tgaangggg agtaggggta tggtattttta 550
W--> 254  tgggaaaana gttttttggc actaaaccnt tttgaattac ctaatanatt 600
W--> 256  tatgtggaaa cctgtccttt ttttncagnt naanaaaaat ttttncccnt 650
W--> 258  gaaantnatt tttagnaagn atatnaaaag nattttttttt ttcaagngtc 700
     260  agaaacctttt tagcatcatt gaagttaaaa tgactgtcca taaactttttc 750
W--> 262  agaaatagta ggcatttnag gcnacnagat ttgtanangg natnttcata 800
W--> 264  gaattatacc agtganttna ccacctgaan cctcttggat cccgtaagca 850
W--> 266  ttctttgcna caaggaaggg aggtatncng ggtaantcct tgaanttttg 900
W--> 268  gacnggaacn atnacttnga atttnannnn nnnnnnnnnn nnnnnngccg 950
W--> 270  cngggncntt tntcgngnn 969
     272  <210> SEQ ID NO: 7
     273  <211> LENGTH: 952
     274  <212> TYPE: DNA
     275  <213> ORGANISM: Homo Sapien
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     279  <222> LOCATION: 1, 3, 7, 23, 30, 38, 49, 53, 64, 68, 84, 90-92, 109, 130, 147,
     280         152, 182, 198, 211, 221-222, 228, 239, 243, 255, 271, 300, 303,
     281         308-309, 314, 322, 331, 337, 340, 345, 359, 377, 381, 385, 412,
     282         422, 452, 463, 469, 475, 541, 547, 586, 777, 780-781, 784-787,
     283         789, 798, 802-803, 825, 831, 838, 840, 853, 855, 863, 865, 872,
     284         885, 889-890, 907, 921-952
     285  <223> OTHER INFORMATION: unknown base
     287  <400> SEQUENCE: 7
W--> 288  ngntttngtt ccctttttttc ccnggtttttn ttttttggnaa aaaaatttna 50
W--> 290  ggnttaaccc aggnaaaanat taaattccaa gggntttggn nngaattccc 100
W--> 292  cggggggttnc ttttaggggt tcctttgacn ttgaaccaaa ggtttcnggc 150
W--> 294  cnggagggggg ggggggaccgg ttttttttccc cngcgtttcc cccgggggntg 200
W--> 296  gggggttgggg ngcccatttg nngaagtnag tggggagggng gantgggaac 250
W--> 298  ccggnagttt tggagaaagg naggttcctt ccttaacccct ggggggttccn 300
W--> 300  ggngcccnng gagnggcagt tngggggaata ntgtttnagn ggttngggggg 350
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RAW SEQUENCE LISTING   DATE: 02/12/2002
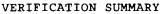PATENT APPLICATION: US/10/052,283   TIME: 12:01:23

Input Set : A:\P2751R1C1.txt
Output Set: N:\CRF3\02122002\J052283.raw

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W--> 302   gttttcctng ggtcccgcca aggggggnggt ncttnataaa agggtgcctt 400
W--> 304   tttccccaca gnttccaggt cngagaggag ccgcaccgtc gggttggaga 450
W--> 306   tngcgcgcaa ggnggcttnt ggttnggatt tgccccgcat cggccacagg 500
W--> 308   aaaagcctgg tccctaggca cggttgtggt tcgagctttt ngttttntcg 550
W--> 310   aacattgagg tattcgctca gcccaccacg ttgtcntcgg ggttattagg 600
     312   ccccagtcac aagccctatg atgttttcag acttcccagg tggagataag 650
     314   gaaaatttta ctatttctgc agaacttctg ttgatgtaca gcattgtatt 700
     316   tagcaacttc tgtgtagatc tgaaaataaa tacattacca attgttagtt 750
W--> 318   gcgttttttat taatataatt ttagagnagn ngannnngnt gttagacnta 800
W--> 320   cnnaggtaaa ttatgtggca ctttngcatt nttgttgntn catgttcccc 850
W--> 322   tgnantttgc ttngngattt cnatttattc caaantcann atagaatgta 900
W--> 324   atttccnaac ccacagtccg nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 950
W--> 326   nn 952
     328   <210> SEQ ID NO: 8
     329   <211> LENGTH: 752
     330   <212> TYPE: DNA
     331   <213> ORGANISM  Homo Sapien
     333   <220> FEATURE:
     334   <221> NAME/KEY  unsure
     335   <222> LOCATION  4-6, 8, 11, 34, 37, 40, 87, 123, 125, 133-134, 138, 143, 147,
     336         166, 171, 174, 199, 242, 263, 302, 305, 333, 363, 380, 423, 425,
     337         463, 527, 534, 539, 547-548, 553, 556, 561, 583, 604, 607, 615,
     338         618, 621, 624, 628, 638, 641-643, 646, 648, 654-655, 658, 660,
     339         666, 674, 676, 679, 688-689, 704, 710-725, 727, 730, 735, 739,
     340         741, 745, 747, 749, 751-752
     341   <223> OTHER INFORMATION: unknown base
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W--> 344   ggannngntt ncaaaatggg atttttaacc aaantanggn agagaaaagt 50
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W--> 348   tttcagagtt caacaggaaa aangngaaca aanngccngg agntttnaaa 150
W--> 350   gttttgggaa agccantttt natntgttca aggaacagtt tttatttgng 200
W--> 352   atgccaatca gaattttgga cccagtataa tcaaggtcag antttcaacc 250
W--> 354   taagcctgga ccngacccat aataacggaa agtttaacaa tgactcacat 300
W--> 356   tntcntaaag tttccagcca gaataggaca cgntcatttg gtcattttcc 350
W--> 358   cggtccagag ttnttggatg tagagaaaan tagcttttcc caggaacaat 400
W--> 360   tttgtgattc cgcaggagaa ggntntgaaa gaatacatca agattttgaa 450
W--> 362   tttggtgatg aanttagcag cagctccact gaacagataa gggcaaccac 500
W--> 364   acctccaaat caaggaaggc cagattntcc tgtntatgnt aaccttnnag 550
W--> 366   aantgnaaat ntcccagtat ggtcttcccc canttcttgg gagcctggta 600
W--> 368   attnagntta ttggngcntg nganactnat atagacanct nnnggngntg 650
W--> 370   ttannatnan cacagnggga catngnatng aagttggnna cctcttgctt 700
W--> 372   ggantcgggn nnnnnnnnnn nnnnncntcn ccgcngggnc ntttntngng 750
W--> 374   nn 752
     376   <210> SEQ ID NO: 9
     377   <211> LENGTH: 478
     378   <212> TYPE: DNA
     379   <213> ORGANISM  Homo Sapien
     381   <400> SEQUENCE  9
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VERIFICATION SUMMARY   DATE: 02/12/2002
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Input Set : **A:\P2751R1C1.txt**
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L:11 M:270 C  Current Application Number differs, Replaced Current Application No
L:11 M:271 C. Current Filing Date differs, Replaced Current Filing Date
L 38 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L 40 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L:46 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:1
L:61 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:2
L:63 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:107 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:109 M 341 W. (46) "n" or "Xaa" used, for SEQ ID#:3
L:111 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L.113 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L·115 M 341 W  (46) "n" or "Xaa" used, for SEQ ID# 3
L:117 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:119 M 341 W. (46) "n" or "Xaa" used, for SEQ ID#:3
L·121 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:3
L:123 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:125 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:3
L:127 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:3
L·129 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:3
L·131 M 341 W. (46) "n" or "Xaa" used, for SEQ ID# 3
L:133 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 3
L:135 M:341 W. (46) "n" or "Xaa" used, for SEQ ID# 3
L:139 M 341 W. (46) "n" or "Xaa" used, for SEQ ID# 3
L.159 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:4
L:161 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:4
L:163 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:4
L:165 M 341 W  (46) "n" or "Xaa" used, for SEQ ID# 4
L:167 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:169 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:171 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L:188 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L.190 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L.192 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:194 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
L:196 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
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L:200 M:341 W  (46) "n" or "Xaa" used, for SEQ ID#:5
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L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:212 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:232 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L:234 M 341 W. (46) "n" or "Xaa" used, for SEQ ID#:6
L:236 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L:238 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 240 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:242 M 341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
```

VERIFICATION SUMMARY
PATENT APPLICATION: **US/10/052,283**

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Output Set: **N:\CRF3\02122002\J052283.raw**

```
L 244 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L 246 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 248 M 341 W  (46) "n" or "Xaa" used, for SEQ ID#:6
L 250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L.17046 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:564
```